Inventors: Hunter and Lu Filing Date: Herewith Attorney Client-Matter No.: 66671-044(P-SC 5712) (858-535-9001)

	·								
	Colony Color pGAD GH Insert								
pAS2 Insert									
	PIN1	PIN2	PIN3						
NIMA	Blue	Blue	Blue						
K40M NIMA	Blue	Blue	Blue						
NIMA280-699	Blue	Blue	White						
NLK1	White	White	Blue						
NIMA280-699FS	White	White	White						

FIG. 1A

NIMA		·	699
	47% identity		
NLK1		445	

FIG. 1B

## TGCTGGCCAGCACCTCGAGGGAAG

				-																	
1	ATG	GCG	GAC	GAG	GAG	AAG	CTG	CCG	CCC	GGC	TGG	GAG	AAG	CGC	ATG	AGC	CGC	AGC	TCA	GGC	
	М	A	D	Ε	Ε	K	Ĺ	Þ	Ď	G	W	Ε	K	R	M	s	P.	S	s	G	20
				' !																	
61 CGAGTGTACTACTTCAACCACATCACTAACGCCAGCCAGTGGGAGCGGCCCAGCGGCAAC																					
01																					
	R	V	Y	<u>Y</u>	<u></u>	N	<u> </u>	I	T	N	_A	S	Q	W	Ε	R	P	S	G	И	40
!																					
121	AGC	AGC	AGT	GGT	GGC	AAA	AAC	GGG	CAG	GGG	GAG	CCT	'GCC	AGG	GTC	CGC	TGC	TCG	CAC	CTG	
	S	S	S	G	G	K	N	G	Q	G	Ε	Þ	A	R	V	R	С	S	н	Ĺ	60
				<u> </u>																	
181	CTG	GTG	AAG	CAC	AGC	CAG	TCA	CGG	CGG	CCC	TCG	TCC	TGG	CGG	CAG	GAG	AAG	ATC	ACC	cgg	1
	}			Н									W				K	I	Т	R	80
	<del>-</del>		<u>-:`</u>				_ <u>~</u> _	<u> </u>	<u> </u>	-	_		, ,	••	~	_	1.	-	•	• `	
																				a. a	100
241				_																.GAG	100
	T	K	Ε	Ε	A	Ļ	Έ	Ŀ	I	N	G	Y	I	Q	K	I	K	S	G	Ε	
										-											٠
301	GAG	GAC	TTT	GAG	TCI	'CTG	GCC	TCA	CAG	TTC	AGC	GAC	TGC	AGC	A	'GCC	AAG	GCC	AGG	GGA	
	E	0	F	Ξ	S	L	A	S	Q	F	\$	D	$\subset$	S	S	A	K	A	R	G	120
												•									
361	GAC	CTG	GGT	GCC	TTC	AGC	AGA	GGT	CAG	ATG	CAG	, A'AC	CCA	TTT	'GA_A	GAC	:GCC	TCG	TTT	'GCG	
	ם	L	G	·A	F	s	R	G	0	М	0	ĸ	p	F	Ε	D	A	S	F	Α.	140
	_		_		-	_	• •	_	~	••	~	•	-	-	_	_	••	·	-	••	
421	CTC									.~~~	mmc							3.00		·cmc	
421	1 -					-														CTC	
	L	R	Т	G	E	<u> </u>	`S	G	_ P		F	T	D	S	G	I	H.	I	I	L ——	160
	ŀ																				
481	CGC	ACT	GAC	#IGA	\GG(	TGC	GGA	GCC	CAC	GCC	TGG	CCI	CGC	GGC	AGC	GC.A	.GGG	CGG	CTA	.GGC	
	R	T.	Ε	*			٠.												<b>.</b> .		•
				_					-										,	-	
541	CGC		\GC1	raco	CCI	TGC			,GCC	AGI	'GGC	:CGA	ACC	ccc	CAC	TCC	CTC	CCA	رحو	TCA	
601																				TGG	
661																				.GGA	
				- (																	
721 ATTGACTTCAGCAGGGGTGGGAGGCTCCCAGACCCAGGGCAGTGTGGGGAGGGGTGTT  781 CCAAAGAGAAGGCCTGGTCAGCAGAGCCGCCCCGTGTCCCCCAGGTGCTGGAGGCAGAC																					
781																					
-841	TCC	AGC	3GC(	IGA.ª	\TTC	GTT	CTA	(GT)	CAGC		\CGC	TCC	TCI	GTT	CAC	TCC	CAA	AGC	TGA	LACA	
901	CTC	ATO	GCGC	GCAC	ICC.	\TGC	GCC	CTC	TGA	NGC A	LAC1	GTC	CAC	ACC	CTI	TCA	CCC		LAT1	AAA	
961	CCC	:AGA	LAC(	CACT	'AA	\AA?	بميم	LAA	LAAA	LAAA								,			

FIG. 2A

P C W E K R M S R S S G R V Y Y F N H I T N A S O W E R P S G N S S S Y T P W T V R Y S K S K K R E Y F F N P E T K H S O W E E P E G T N K D A G W E M A K T S S - G Q R Y F L N H I D Q T T T W Q D P R K A M L S P G W E E R Q D V L - G R T Y Y V N H E S R R T Q W K R P S P D D D L P G W E R R T D N F - G R T Y Y V D H N T R T T T W K R P T L D Q T E G P W E R A I S P N K V P - Y Y I N H E T Q T T C W D H P K M T E L Y A G W M R V Q D T S - G - T Y Y W H I P T G T T Q W E P P G R A S P S	Р. СИЕ G. ХУ. N Н. Т. Т. И. Р	K H S Q S R R P S S W R Q E K I T R T K E E A L E L I N G Y K H K D S R R P A S H R S E N I T I S K Q D A T D E L K T L	- EDFESLASQFSDCSSAKA-RGDLGAFSR-GQMQTNSFEALAKERSDCSSYKR-GGDLGAFSR-GQMQ-ADFGKLAKKHSTCPSGKR-GGDLGEF-RQGQMVFEDLAKEYSTDSSASK-GGDLGWFAKEGQMDDAKFSELAKEKSTDPGSKNQGGELGWF-DQSTMV-ANFEEVATRVSDCSSAKR-GGDLGSFGR-GQMQ	FE.LAKS.CSS.KGGDLG.F.R.G	A S, F A L R T G E M S G - P V F T D S G I H I I L R T E 164 Pinl/Human A A F Q L K V G E V S D - I V E S G S G V H V I K R V G 170 ESS1/SC V V F S C P V L E P T G - P L H T Q F G Y H I I K V L Y 93 Parvulin/EC A A F K L K T G E V S D - P V K T Q Y G Y H I I K K T E 215 PrsA/BS A A F A L K N G T I T T T P V K T N F G Y H I I K K T E 215 Cbf2/CJ A T Y A L K V G D I S D - I V D T D S G V H I I K R T E 69 EST/AT A A F A F L K C G E S P V T G Y H I I K Consensus
X X P P B C X C C C C C C C C C C C C C C C C C		H I I I I I I I I I I I I I I I I I I I	X X X X X X X X X X X X X X X X X X X	· · · ·	7 7 7 7 7 8 9 9 9 9 9 9 9 9 9 9 9 9 9 9
Pin1,/Human         5         E           ESS1/9C         9         T           Yap/Human         171         V           Nectd/ // // // // // // // // // // // // /	Consensus	Pin1/Human 59 H ESS1/SC 63 H Parvulin/EC 8 H PrsA/BS 140 H Cbf2/CJ 137 H EST/AT			FIG. 2C



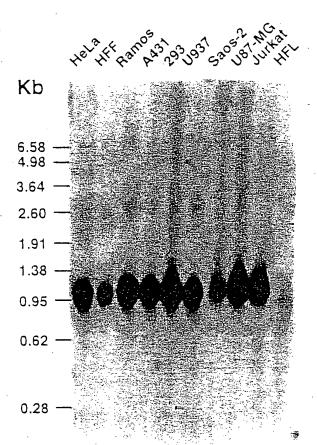
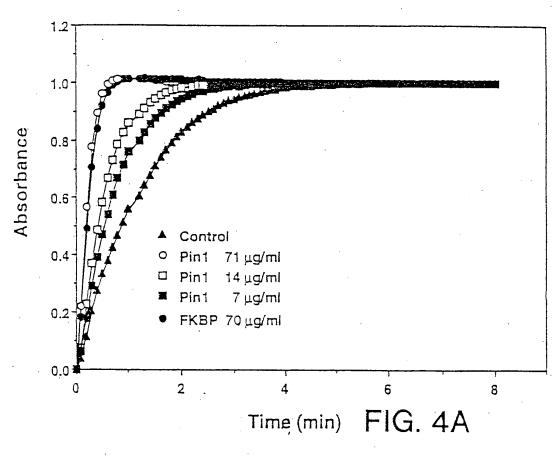
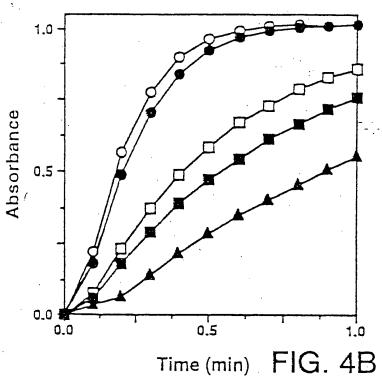
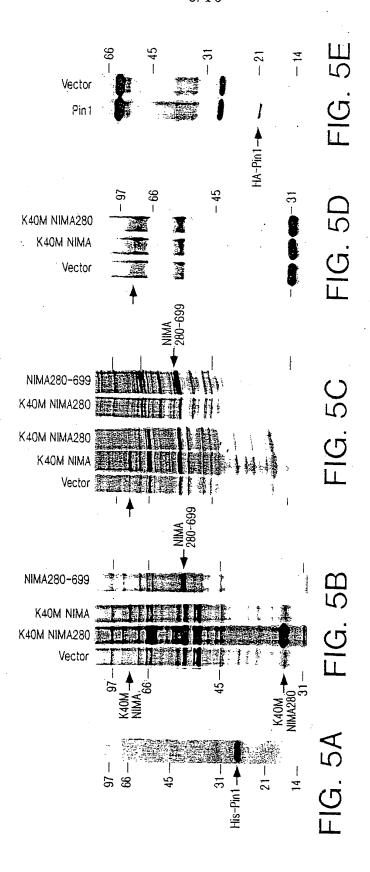


FIG. 3







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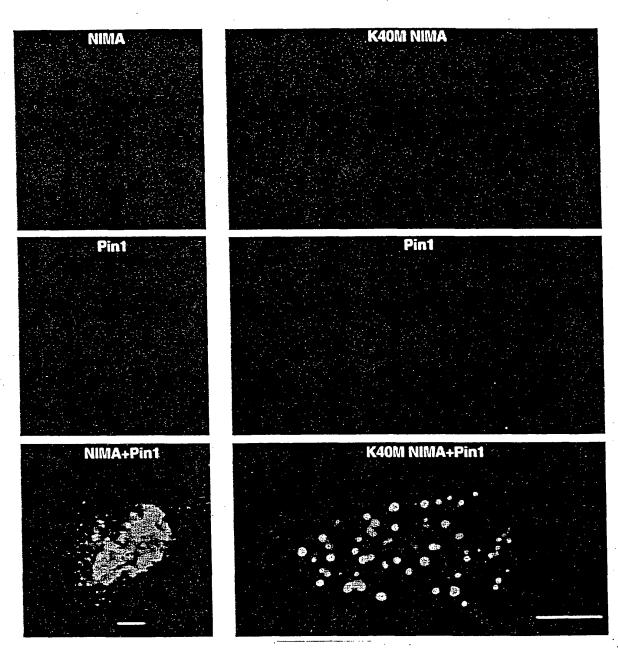


FIG. 6

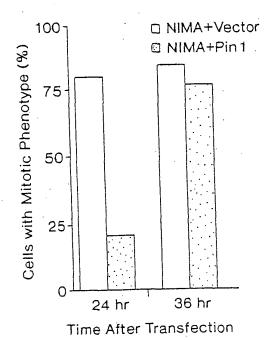
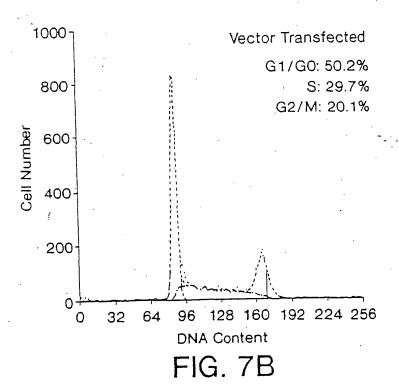


FIG. 7A



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